

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: May 4, 2005, 12:06:08 ; Search time 2533.16
Seconds
(without alignments)
5092.486 Million cell
updates/sec

Title: US-09-300-482-225
Perfect score: 316
Sequence: 1
gataagggtgcgcacactgag.....gaatggagggagtacttcga 316

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 45554873 seqs, 20411521753 residues

Total number of hits satisfying chosen parameters: 91109746

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_NA_Main:
1: /cgn2_6/ptodata/1/pna/PCTUS1_COMB.seq:/*
2: /cgn2_6/ptodata/1/pna/PCTUS2_COMB.seq:/*
3: /cgn2_6/ptodata/1/pna/PCTUS_COMB.seq:/*
4: /cgn2_6/ptodata/1/pna/US06_COMB.seq:/*
5: /cgn2_6/ptodata/1/pna/US07_COMB.seq:/*
6: /cgn2_6/ptodata/1/pna/US080_COMB.seq:/*
7: /cgn2_6/ptodata/1/pna/US081_COMB.seq:/*
8: /cgn2_6/ptodata/1/pna/US082_COMB.seq:/*
9: /cgn2_6/ptodata/1/pna/US083_COMB.seq:/*
10: /cgn2_6/ptodata/1/pna/US084_COMB.seq:/*
11: /cgn2_6/ptodata/1/pna/US085_COMB.seq:/*
12: /cgn2_6/ptodata/1/pna/US086_COMB.seq:/*
13: /cgn2_6/ptodata/1/pna/US087_COMB.seq:/*
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